



Collection, Concentration and Long Term Room Temperature Stabilization of Forensic DNA in Liquid Format is a Reality

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1. Introduction

Crime scene investigators are trained to air-dry evidentiary samples following swab-based collection. Drying swabs can lead to irreversible binding of DNA to the swab material resulting in low DNA recovery. However this step can be challenging, especially in areas of high humidity and air moisture content. There are several factors affecting DNA integrity and yields of evidentiary samples:

- ☐ Irreversible binding to swab material
- □ Nucleases
- ☐ Microbial contamination
- □ Temperature fluctuations
- ☐ Heat
- ☐ Humidity
- UV light
- ☐ Improper collection
- ☐ Storage conditions and length of backlog
- ☐ Transport conditions & transit time

Mawi has developed iSWAB-ID, an efficient liquid-based sample collection system which utilizes swabs. iSWAB-ID enables long term room temperature stabilization of the collected sample at the point of collection while ensuring proper chain of custody. This system maximizes sample recovery and simplifies sample processing extensively over current practices, allowing for Enhanced ID Profiling.

Evidentiary Material Processing Bottlenecks



2. Objectives

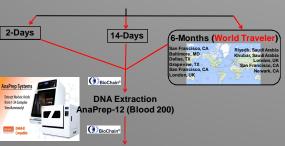
- To assess the efficiency of ISWAB-ID in the collection, concentration, and long-term room temperature stabilization of both reference and evidentiary samples by assessing the usability of purified DNA in human ID-STR profiling assays.
- To assess the stability of ISWAB-ID samples during multistop-global transport process

3. Experimental Design

Collect reference and mocked evidentiary samples into the iSWAB-ID Sample Collection Device (400 µL)

Transport to processing lab by standard mail Store samples at room temperature

100 µL aliquoted from each iSWAB-ID sample post collection



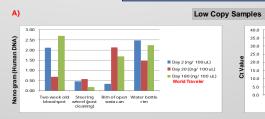
Quantify DNA (QIAGEN Investigator Quantiplex)

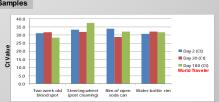


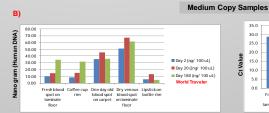
STR profiling (Promega PowerPlex 16 HS) and analysis

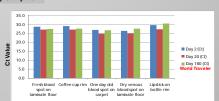
4. Results

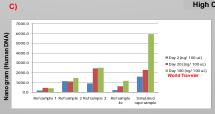
I- DNA Extraction and Human DNA Quantification iSWAB-ID Efficiently Recovers and Stabilizes Human DNA from Both Reference & Evidentiary Samples in Long Term, Room Temperature Storage from Low, Medium and High Copy Number Samples

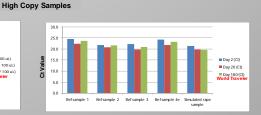












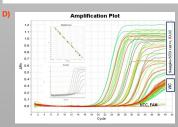
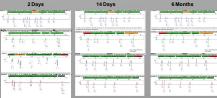


Fig 1. A selection of real-world, reference, and simulated evidence samples was collected in iSWAB-ID devices and stored at room temperature. DNA was extracted from 100 µL aliquots, using the AnaPrep Blood DNA extraction kit on AnaPrep 13 instrument, either two days or two weeks after collection. In all cases, PCR-amplifiable Human DNA was recovered from stabilized samples after storage for two weeks at room temperature. All samples were quantified by QIAGEN Investigator Quantiplex Kit targeting Human DNA. A) Low copy samples, B) Medium copy samples, C) High copy samples, D) Amplification plot for all samples including standard curve, VIC, (two & 14 days and 6 months samples), ND: Not Detected

II- STR Profiling and Analysis with Promega PowerPlex 16 HS First Pass STR Profiling Analysis Samples are Suitable for Comparison Purposes

Lipstick on a Cup Rim: An Example of Touch DNA Sample





5. Summary & Conclusions

Evidentiary & Reference Samples Processing • The iSWAB-ID sample collection device efficiently recovered and stabilized DNA of forensic significance in liquid Bottlenecks: All Resolved with iSWAB-ID format at the point of collection. cted and stabilized in iSWAB-ID at ambient 1 2 3 4 5 6 7 8 ure remained of sufficient quality to analyze for at

Ŧ	Description	Time Spent (mins)		
	Screen Identify Collect		Control of the Contro	 DNA col temperatu
	No Drying Required	0 min/Sample	4	least 6 mg
	Package			
	Reporting and documentation			
	Transport			Unlike :
	Store			
	Cut swab/ Punch FTA	0 min/Sample	9	with iSW
	Card	Not Applicable		runs and
	Lyse			
ī	Extract			 Extende
	Quantify			- Exteriue
	STR profiling	5-10% Further Analysis	13	space-sav
	Data Analysis			voordting of

Significant Improvement on Processing Efficiency ✓ Faster Collection ✓ Higher First Pass Rate
✓ Higher Sample processing Throughput

- samples processed from swabs, samples collected /AB-ID yield more DNA allowing multiple analytical d sufficient material for archiving.
- ving storage and eliminates sample degradation from excessive backlogs.

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